

FIGURE 1A

ATGGAGAGCAAGGTGCTGCTGGCCGTCGCCCTGTGGCTCTGCGTGGAGACCC
 GGGCCGCCTCTGTGGGTTTGCCTAGTGTCTCTTGATCTGCCCAGGCTCAGCA
 TACAAAAAGACATACTTACAATTAAGGCTAATACAACTCTTCAAATTACTTGCAG
 GGGACAGAGGGGACTTGGACTGGCTTTGGCCCAATAATCAGAGTGGCAGTGAG
 CAAAGGGTGGAGGTGACTGAGTGCAGCGATGGCCTCTTCTGTAAGACACTCAC
 AATTCCAAAAGTGATCGGAAATGACACTGGAGCCTACAAGTGCTTCTACCGGG
 AAAGTGAAGTGGCCTCGGTCAATTTATGTCTATGTTCAAGATTACAGATCTCCATT
 TATTGCTTCTGTTAGTGACCAACATGGAGTCGTGTACATTACTGAGAACAAAA
 CAAAAGTGTGGTGATTCCATGTCTCGGGTCCATTTCAAATCTCAACGTGTCACTT
 TGTGCAAGATACCCAGAAAAAGAGATTTGTTCTGATGGTAACAGAAATTTCTGG
 GACAGCAAGAAGGGCTTTACTATTCCAGCTAGATGATCAGCTATGCTGGCATG
 GTCTTCTGTGAAGCAAAAATTAATGATGAAAGTTACCAGTCTATTATGTACATAG
 TTGTGCTTGTAGGGTATAGGATTTATGATGTGGTTCTGAGTCCGTCTCATGGAA
 TTGAAGTATCTGTTGGAGAAAAGCTTGTCTTAAATTGTACAGCAAGAACTGAAC
 TAAATGTGGGGATTGACTTCAACTGGGAATACCCCTTCTTGAAGCATCAGCATA
 AGAAACTTGTAAACCGAGACCTAAAAACCCAGTCTGGGAGTGAGATGAAGAAA
 TTTTGTGACACCTTAACTATAGATGGTGTAAACCCGGAGTGACCAAGGATTGTAC
 ACCTGTGCAGCATCCAGTGGGCTGATGACCAAGAAGAACAGCACATTTGTGAG
 GGTCCATGAAAAACCTTTTGTGCTTTTGGAAAGTGGCATGGAATCTCTGGTGGA
 AGCCACGGTGGGGGAGCGTGTGAGAATCCCTGCGAAGTACCTTGGTTACCCAC
 CCCAGAAATAAAATGGTATAAAATGGAATAACCCCTTGAAGTCCAATCACACA
 TTAAGCGGGGATGTAAGTACGATTATGGAAGTGAGTGAAAGAGACACAGGA
 AATTACACTGTCATCTTACCAATCCCATTTCAAAGGAGAAGCAGAGCCATGTG
 GTCTCTCTGGTTGTGTATGTCCACCCAGATTGGTGAGAAATCTCTAATCTCTC
 CTGTGGATTCTACAGTACGGCACCACTCAAACGCTGACATGTACGGTCTATG
 CCATTCTCCCCCGCATCACATCCACTGGTATTGGCAGTTGGAGGAAGAGTGC
 GCCAACGAGCCCAGCCAAGCTGTCTCAGTGACAAACCCATACCCCTTGTGAAGA
 ATGGAGAAGTGTGGAGGACTTCCAGGGAGGAAATAAAATTGAAGTTAATAAAA
 ATCAATTTGCTCTAATTGAAGGAAAAAACAAACTGTAAAGTACCCCTTGTATCCA
 AGCGGCAAATGTGTCAGCTTTGTACAAATGTGAAGCGGTCAACAAAGTCGGGA
 GAGGAGAGAGGGTGATCTCCTTCCACGTGACCAGGGGTCTGAAATTAATTTG
 CAACCTGACATGCAGCCCACTGAGCAGGAGAGCGTGTCTTTGTGGTGCACCTGC
 AGACAGATCTACGTTTGAGAACCTCACATGGTACAAGCTTGGCCACAGCCTCT
 GCCAATCCATGTGGGAGAGTTGCCACACCTGTTTGCAAGAACTTGGATACTCT
 TTGGAAATTGAATGCCACCATGTTCTCTAATAGCACAAATGACATTTTGATCATG
 GAGCTTAAGAATGCATCCTTGCAGGACCAAGGAGACTATGTCTGCCTTGTCAA
 GACAGGAAGAAGCAAGAAAGACATTGCGTGGTTCAGGCAGCTCACAGTCCTAGA
 GCGTGTGGCACCCACGATCACAGGAAACCTGGAGAATCAGACGACAAGTATTG
 GGGAAAGCATCGAAGTCTCATGCACGGCATCTGGGAATCCCCCTCCACAGATC
 ATGTGGTTTAAAGATAATGAGACCTTGTAGAAGACTCAGGCATTGTATTGAAG
 GATGGGAACCGGAACCTCACTATCCGCAGAGTGAGGAAGGAGGACGAAGGCC
 TCTACACCTGCCAGGCATGCAGTGTCTTGGCTGTGCAAAAGTGGAGGCATTTT
 TCATAATAGAAGGTGCCAGGAAAAGACGAACCTTGGAAATCATTATTCTAGTAG
 GCACGGCGGTGATTGCCATGTTCTTCTGGCTACTTCTTGTTCATCATCCTACGGA
 CCGTTAAGCGGGCCAATGGAGGGGAACCTGAAGACAGGGTACCTGTCCATCGT
 CATGGACCCAGATGAACTCCCATTTGGATGAACATTGTGAACGACTGCCTTATGA
 TGCCAGCAAATGGGAATTTCCACAGAGACCGGCTGAAGCTAGGTAAGCCTCTTG
 GCCGTGGTGCCTTTGGCCAAGTGATTGAAGCAGATGCCTTTGGAATTGACAAG
 ACAGCAACTTGCAGGACAGTAGCAGTCAAAATGTTGAAAGAAGGAGCAACACA
 CAGTGAGCATCGAGCTCTCATGTCTGAACTCAAGATCCTCATTATGTTGGTCA
 CCATCTCAATGTGGTCAACCTTCTAAGTGCCTGTACCAAGCCAGGAGGGCCAC
 TCAAGGTGATTGTGAATTCTGCAAAATTTGGAAACCTGTCCACTTACCTGAGGA
 GCAAGAGAAATGAATTTGTCCCTACAAGACCAAGGGGCACGATTCCGTCAA
 GGGAAAGACTACGTTGGAGCAATCCCTGTGGATCTGAAACGGCGCTTGGACAG

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FIGURE 1B

CATCACCAGTAGCCAGAGCTCAGCCAGCTCTGGATTTGTGGAGGAGAAGTCCC
TCAGTGATGTAGAAGAAGAGGAAGCTCCTGAAGATCTGTATAAGGACTTCCTG
ACCTTGGAGCATCTCATCTGTTACAGCTTCCAAGTGGCTAAGGGCATGGAGTTC
TTGGCATCGCGAAAGTGTATCCACAGGGACCTGGCGGCACGAAATATCCTCTT
ATCGGAGAAGAACGTGGTTAAAATCTGTGACTTTGGCTTGGCCCGGGATATTTA
TAAAGATCCAGATTATGTCAGAAAAGGAGATGCTCGCCTCCCTTTGAAATGGAT
GGCCCCAGAAACAATTTTGTGACAGAGTGTACACAATCCAGAGTGACGTCTGGT
CTTTTGGTGTGTTTGTGCTGTGGGAAATATTTTCTTAGGTGCTTCTCCATATCCTGG
GGTAAAGATTGATGAAGAATTTTGTAGGCGATTGAAAGAAGGAACTAGAATGA
GGGCCCCCTGATTATACTACACCAGAAATGTACCAGACCATGCTGGACTGCTGG
CACGGGGAGCCCAGTCAGAGACCCACGTTTTTCAGAGTTGGTGGAAACATTTGGG
AAATCTCTTGCAAGCTAATGCTCAGCAGGATGGCAAAGACTACATTGTTCTTCC
GATATCAGAGACTTTGAGCATGGAAGAGGATTCTGGACTCTCTCTGCCTACCTC
ACCTGTTTCTGTATGGAGGAGGAGGAAGTATGTGACCCCAAATTCATTATGA
CAACACAGCAGGAATCAGTCAGTATCTGCAGAACAGTAAGCGAAAGAGCCGGC
CTGTGAGTGTAAAAACATTTGAAGATATCCCGTTAGAAGAACCAGAAGTAAAAG
TAATCCCAGATGACAACCAGACGGACAGTGGTATGGTTCCTTGCCTCAGAAGAG
CTGAAAACCTTTGGAAGACAGAACCAAATTATCTCCATCTTTTGGTGGAAATGGT
CCCAGCAAAGCAGGGAGTCTGTGGCATCTGAAGGCTCAAACCAGACAAGCG
GCTACCAGTCCGGATATCACTCCGATGACACAGACACCACCGTGTACTCCAGT
GAGGAAGCAGAACTTTTAAAGCTGATAGAGATTGGAGTGCAAACCGGTAGCAC
AGCCCAGATTCTCCAGCCTGACTCGGGGACCACACTGAGCTCTCCTCCTGTTTA
A

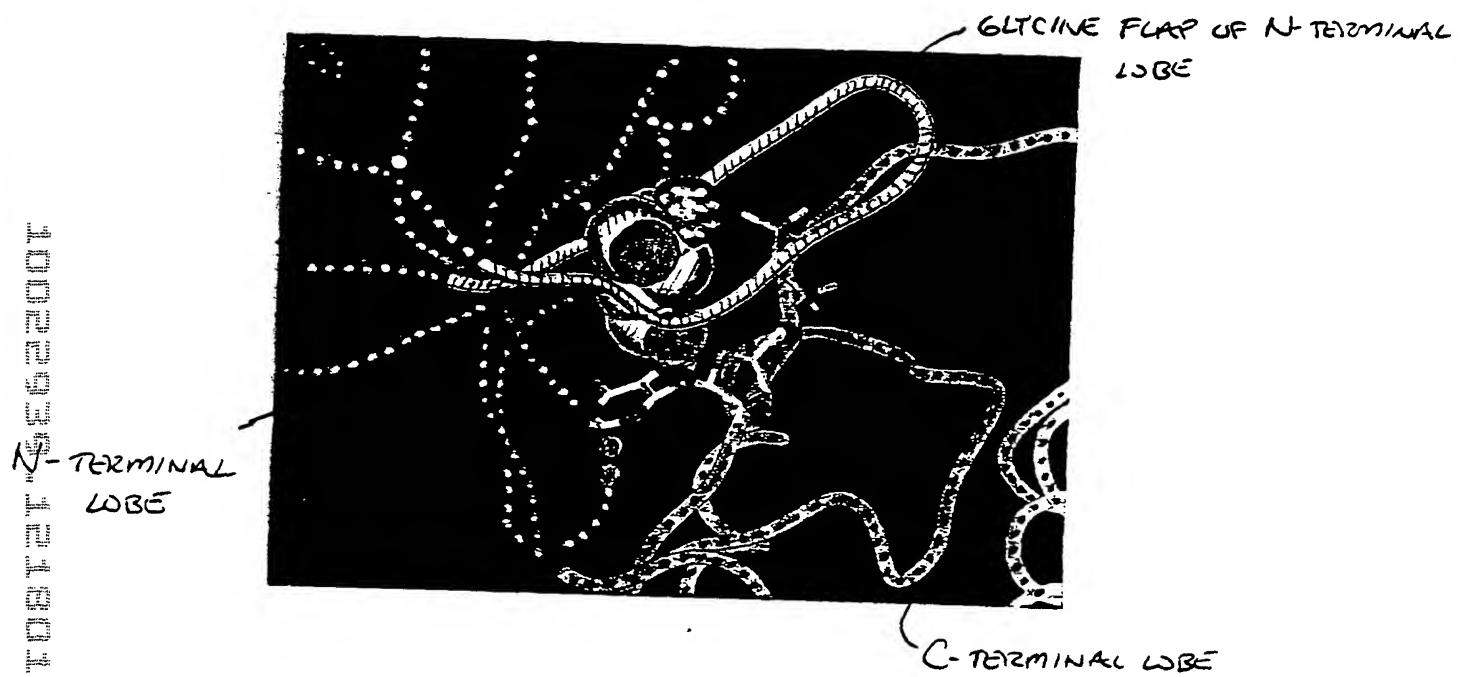
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FIGURE 2

MESKVLLAVALWLCVETRAASVGLPSVSLDLPRLSIQKDILTIKANTTLQITCRGQR
DLDWLWPNNQSGSEQRVEVTECSDGLFCKTLTPKVIGNDTGAYKCFYRETDLAS
VTYVYVQDYRSPFLASVSDQHGVVYTENKNKTVVIPCLGSSISNLNVSLCARYPEKR
FVPDGNRISWDSKKGFTIPSYMISYAGMVFCFAKINDESYQSIMYTVVVVGYRIYDV
VLSPSHGIELSVGEKLVNCTARTELNVGIDFNWEYPSSKHQHKKLVNRDLKTQS
GSEMKKFLSTLTIDGVTRSDQGLYTCAASSGLMTKKNSTFVRVHEKPFVAFGSGM
ESLVEATVGERVRIPAKYLGYPPEIKWYKNGIPLESNHTIKAGHVLTMEVSESDT
GNYTVILTNPISKEKQSHVVSLVYVPPQIGEKSLISPVDSYQYGTQTTLCTVYAI
PPHHIHWYWQLEEECANEPSQAVSVTNYPCEEWRVSVDFQGGNKIEVNKNQFA
LIEGKNKTVSTLVIQAANVSALYKCEAVNKVGRGERVISFHVTRGPEITLQPDMP
TEQESVSLWCTADRSTFENLTWYKLGPOPLPHVGEPLTPVCKNLDTLWKLNATM
FSNSTNDILMELKNASLQDQGDYVCLAQDRKTKKRHCVRQLTVLERVAPTITGN
LENQTTSIGESIEVSCTASGNPPPQIMWFKDNETLVEDSGIVLKDGNRNLTIRRVK
EDEGLYTCQACSVLGCAKVEAFFIEGAQEKTNLEIILVGTAVIAMFFWLLLVIILRT
VKRANGGELKTGYLSIVMDPDELPLDEHCERLPYDASKWEFPRDRLKLGKPLGRG
AFGQVIEADAFGIDKTATCRTVAVKMLKEGATHSEHRALMSELKILIHGHILNVV
NLLGACTKPGGPLMVTVEFCKFGNLSTYLRSKRNEFVPYKTKGARFRQGDYVG
AIPVDLKRRLDSITSSQSSASSGFVEEKSLSDEVVEEEAPEDLYKDFTLEHLICYSFQ
VAKGMEFLASRKCIHRDLAARNILLSEKNVVKICDFGLARDIYKDPDYVRKGDAR
LPLKWMAPETIFDRVYTIQSDVWSFGVLLWEIFSLGASPYPGVKIDEEFCRRLKEGT
RMRAPDYTTPEMYQTMLD CWHGEPSQRPTFSELVEHLGNLLQANAQQDGKDYTVL
PISETLSMEEDSGLSLPTSPVSCMEEEVCDPKFHYDNTAGISQYLQNSKRKSRPVS
VKTFEDIPLEPEVKVIPDDNQTDSGMVLASEELKTLEDRTKLSPSFGGMVPSKSRE
SVASEGSNQTSQGYQSGYHSDDTDTTVYSSEEAEILLKLEIGVQTGSTAQILQPDSGT
TLSSPPV

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FIGURE 3A



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FIGURE 3B

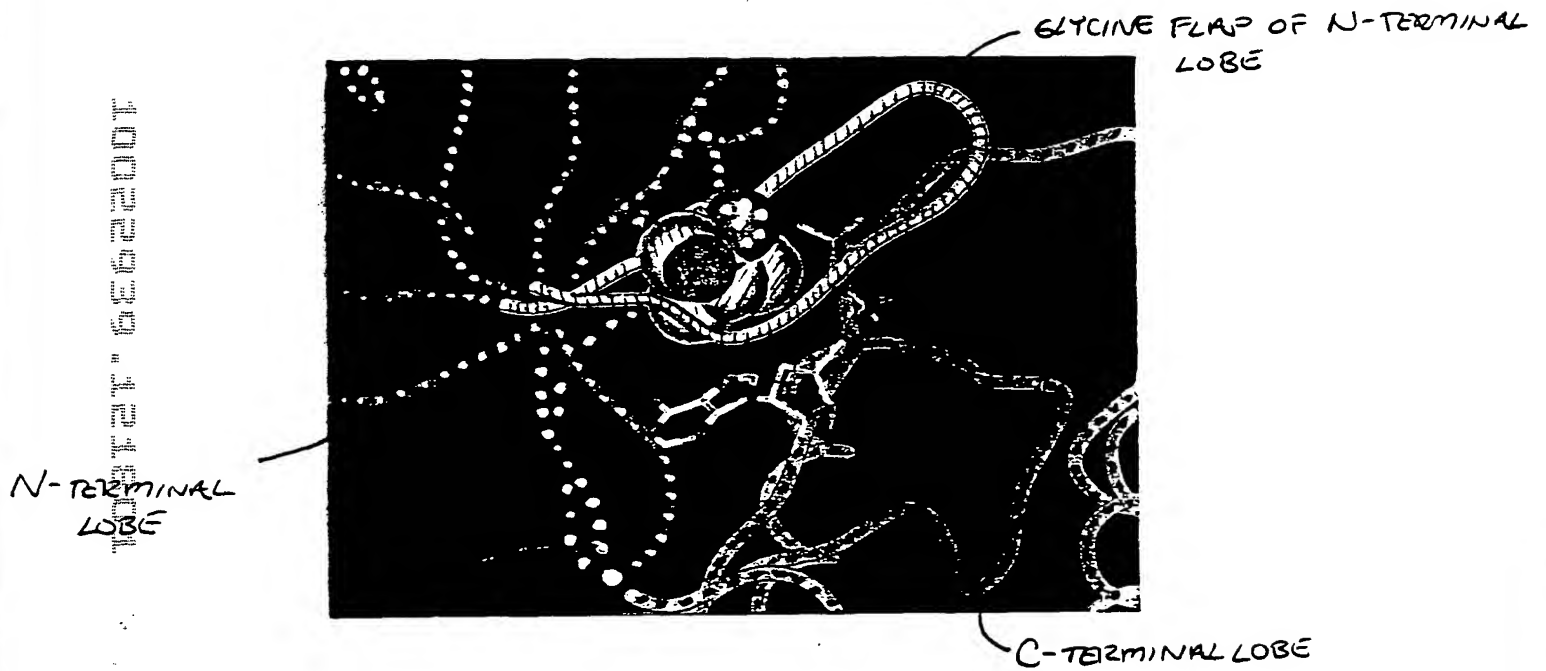


FIGURE 4A

Anti-phosphotyrosine

E848 V848

12 12 120 12
- + + - - +

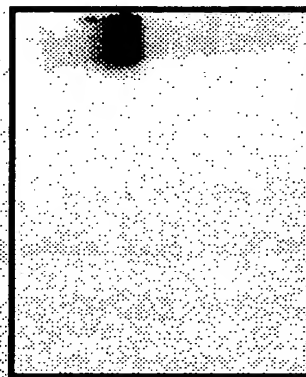


FIGURE 4B

Anti-KDR

E848 V848

120 12
- -
Enzyme (ng)
ATP (1 mM)

